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Sequence Listing

<110> Hofmann, Kay

<120> Protease

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<140> US 09/869,309

<141> 2001-07-20

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<151> 1999-01-22

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<151> 1999-06-08

<150> 19929115.2

<151> 1999-06-24

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<170> PatentIn Ver. 2.1

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<211> 592

<212> PRT

<213> Homo sapiens

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 35 40 45

Asn Pro Gln Trp Ala His Leu Pro His Asp Leu Ser Lys Ala Ser Phe
 50 55 60

Leu Gln Leu Arg Asn Trp Thr Ala Ser Leu Leu Cys Ser Ala Ala Asp
 65 70 75 80

Leu Pro Ala Arg Gly Phe Ser Asn Gln Ile Pro Leu Val Ala Arg Gly
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Asn Cys Thr Phe Tyr Glu Lys Val Arg Leu Ala Gln Gly Ser Gly Ala
 100 105 110

Arg Gly Leu Leu Ile Val Ser Arg Glu Arg Leu Val Pro Pro Gly Gly
 115 120 125

Asn Lys Thr Gln Tyr Asp Glu Ile Gly Ile Pro Val Ala Leu Leu Ser
 130 135 140

Tyr Lys Asp Met Leu Asp Ile Phe Thr Arg Phe Gly Arg Thr Val Arg
 145 150 155 160

Ala Ala Leu Tyr Ala Pro Lys Glu Pro Val Leu Asp Tyr Asn Met Val
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Ile Ile Phe Ile Met Ala Val Gly Thr Val Ala Ile Gly Gly Tyr Trp
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Ala Gly Ser Arg Asp Val Lys Lys Arg Tyr Met Lys His Lys Arg Asp
 195 200 205
 Asp Gly Pro Glu Lys Gln Glu Asp Glu Ala Val Asp Val Thr Pro Val
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 Met Thr Cys Val Phe Val Val Met Cys Cys Ser Met Leu Val Leu Leu
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 Tyr Tyr Phe Tyr Asp Leu Leu Val Tyr Val Val Ile Gly Ile Phe Cys
 245 250 255
 Leu Ala Ser Ala Thr Gly Leu Tyr Ser Cys Leu Ala Pro Cys Val Arg
 260 265 270
 Arg Leu Pro Phe Gly Lys Cys Arg Ile Pro Asn Asn Ser Leu Pro Tyr
 275 280 285
 Phe His Lys Arg Pro Gln Ala Arg Met Leu Leu Leu Ala Leu Phe Cys
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 Val Ala Val Ser Val Val Trp Gly Val Phe Arg Asn Glu Asp Gln Trp
 305 310 315 320
 Ala Trp Val Leu Gln Asp Ala Leu Gly Ile Ala Phe Cys Leu Tyr Met
 325 330 335
 Leu Lys Thr Ile Arg Leu Pro Thr Phe Lys Ala Cys Thr Leu Leu Leu
 340 345 350
 Leu Val Leu Phe Leu Tyr Asp Ile Phe Phe Val Phe Ile Thr Pro Phe
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 Leu Thr Lys Ser Gly Ser Ser Ile Met Val Glu Val Ala Thr Gly Pro
 370 375 380
 Ser Asp Ser Ala Thr Arg Glu Lys Leu Pro Met Val Leu Lys Val Pro
 385 390 395 400
 Arg Leu Asn Ser Ser Pro Leu Ala Leu Cys Asp Arg Pro Phe Ser Leu
 405 410 415
 Leu Gly Phe Gly Asp Ile Leu Val Pro Gly Leu Leu Val Ala Tyr Cys
 420 425 430
 His Arg Phe Asp Ile Gln Val Gln Ser Ser Arg Val Tyr Phe Val Ala
 435 440 445
 Cys Thr Ile Ala Tyr Gly Val Gly Leu Leu Val Thr Phe Val Ala Leu
 450 455 460
 Ala Leu Met Gln Arg Gly Gln Pro Ala Leu Leu Tyr Leu Val Pro Cys
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 485 490 495
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 515 520 525
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 530 535 540

Trp Pro Ala Glu Gln Ser Pro Lys Ser Arg Thr Ser Glu Glu Met Gly
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<213> Homo sapiens

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Trp Thr Ala Leu Pro Ser Thr Leu Glu Asn Ala Thr Ser Ile Ser Leu
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Met Asn Leu Thr Ser Thr Pro Leu Cys Asn Leu Ser Asp Ile Pro Pro
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Val Gly Ile Lys Ser Lys Ala Val Val Val Pro Trp Gly Ser Cys His
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Phe Leu Glu Lys Ala Arg Ile Ala Gln Lys Gly Gly Ala Glu Ala Met
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Glu Phe Pro Asp Val Lys Ile Leu Ile Ala Phe Ile Ser Tyr Lys Asp
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Phe Arg Asp Met Asn Gln Thr Leu Gly Asp Asn Ile Thr Val Lys Met
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Tyr Ser Pro Ser Trp Pro Asn Phe Asp Tyr Thr Met Val Val Ile Phe
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Val Ile Ala Val Phe Thr Val Ala Leu Gly Gly Tyr Trp Ser Gly Leu
180 185 190

Val Glu Leu Glu Asn Leu Lys Ala Val Thr Thr Glu Asp Arg Glu Met
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Arg Lys Lys Lys Glu Glu Tyr Leu Thr Phe Ser Pro Leu Thr Val Val
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 Asp Met Pro Glu Thr Ile Thr Ser Arg Asp Ala Ala Arg Phe Pro Ile
 65 70 75 80
 Ile Ala Ser Cys Thr Leu Leu Gly Leu Tyr Leu Phe Phe Lys Ile Phe
 85 90 95
 Ser Gln Glu Tyr Ile Asn Leu Leu Leu Ser Met Tyr Phe Phe Val Leu
 100 105 110
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 165 170 175
 Leu Leu Arg Lys His Trp Ile Ala Asn Asn Leu Phe Gly Leu Ala Phe
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 Ser Leu Asn Gly Val Glu Leu Leu His Leu Asn Asn Val Ser Thr Gly
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 Cys Ile Leu Leu Gly Gly Leu Phe Ile Tyr Asp Val Phe Trp Val Phe
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 Gly Thr Asn Val Met Val Thr Val Ala Lys Ser Phe Glu Ala Pro Ile
 225 230 235 240
 Lys Leu Val Phe Pro Gln Asp Leu Leu Glu Lys Gly Leu Glu Ala Asn
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 Asn Phe Ala Met Leu Gly Leu Gly Asp Val Val Ile Pro Gly Ile Phe
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 Ile Ala Leu Leu Leu Arg Phe Asp Ile Ser Leu Lys Lys Asn Thr His
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 Thr Tyr Phe Tyr Thr Ser Phe Ala Ala Tyr Ile Phe Gly Leu Gly Leu
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 Thr Ile Phe Ile Met His Ile Phe Lys His Ala Gln Pro Ala Leu Leu
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 Tyr Leu Val Pro Ala Cys Ile Gly Phe Pro Val Leu Val Ala Leu Ala
 325 330 335
 Lys Gly Glu Val Thr Glu Met Phe Ser Tyr Glu Glu Ser Asn Pro Lys
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 <213> Homo sapiens

<400> 4

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Ser Val Ser Leu Leu Val Met Phe Phe Phe Phe Asp Ser Val Gln Val
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Val Phe Thr Ile Cys Thr Ala Val Leu Ala Thr Ile Ala Phe Ala Phe
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Leu Leu Leu Pro Met Cys Gln Tyr Leu Thr Arg Pro Cys Ser Pro Gln
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Asn Lys Ile Ser Phe Gly Cys Cys Gly Arg Phe Thr Ala Ala Glu Leu
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Met Ile Ala Phe Val Arg Leu Pro Ser Leu Lys Val Ser Cys Leu Leu
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 195             200             205

Tyr Ile Phe Asn Ser Asn Val Met Val Lys Val Ala Thr Gln Pro Ala
 210             215             220

Asp Asn Pro Leu Asp Val Leu Ser Arg Lys Leu His Leu Gly Pro Asn
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Val Gly Arg Asp Val Pro Arg Leu Ser Leu Pro Gly Lys Leu Val Phe
 245             250             255

Pro Ser Ser Thr Gly Ser His Phe Ser Met Leu Gly Ile Gly Asp Ile
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 <213> Mus musculus

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 35 40 45
 Met Val Met Lys Thr Gly Gln Pro Ala Leu Leu Tyr Leu Val Pro Cys
 50 55 60
 Thr Leu Ile Thr Val Ser Val Val Ala Trp Ser Arg Lys Glu Met Lys
 65 70 75 80
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Phe Gly Ala Leu Pro Ser Val Arg Cys Ala Arg Gly Lys Ser Ser Ser
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Asp Met Pro Glu Thr Ile Thr Ser Arg Asp Ala Ala Arg Phe Pro Ile
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Gly Ile Leu Ala Leu Ser His Thr Ile Ser Pro Phe Met Asn Lys Phe
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Phe Pro Ala Asn Phe Pro Asn Arg Gln Tyr Gln Leu Leu Phe Thr Gln
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Gly Ser Gly Glu Asn Lys Glu Glu Ile Ile Asn Tyr Glu Phe Asp Thr
145 150 155 160
Lys Asp Leu Val Cys Leu Gly Leu Ser Ser Val Val Gly Val Trp Tyr
165 170 175
Leu Leu Arg Lys His Trp Ile Ala Asn Asn Leu Phe Gly Leu Ala Phe
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195 200 205
Cys Ile Leu Leu Gly Gly Leu Phe Ile Tyr Asp Ile Phe Trp Val Phe
210 215 220
Gly Thr Asn Val Met Val Thr Val Ala Lys Ser Phe Glu Ala Pro Ile
225 230 235 240
Lys Leu Val Phe Pro Gln Asp Leu Leu Glu Lys Gly Leu Glu Ala Asp
245 250 255
Asn Phe Ala Met Leu Gly Leu Gly Asp Ile Val Ile Pro Gly Ile Phe
260 265 270
Ile Ala Leu Leu Leu Arg Phe Asp Ile Ser Leu Lys Lys Asn Thr His
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Thr Tyr Phe Tyr Thr Ser Phe Ala Ala Tyr Ile Phe Gly Leu Gly Leu
290 295 300
Thr Ile Phe Ile Met His Ile Phe Lys His Ala Gln Pro Ala Leu Leu
305 310 315 320
Tyr Leu Val Pro Ala Cys Ile Gly Phe Pro Val Leu Val Ala Leu Ala
325 330 335
Lys Gly Glu Val Ala Glu Met Phe Ser Tyr Glu Glu Ser Asn Pro Lys
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 Thr Gly His Trp Leu Leu Met Asp Ala Leu Ala Met Gly Leu Cys Val
 35 40 45
 Ala Met Ile Ala Phe Val Arg Leu Pro Ser Leu Lys Val Ser Cys Leu
 50 55 60
 Leu Leu Ser Gly Leu Leu Ile Tyr Asp Val Phe Trp Val Phe Phe Ser
 65 70 75 80
 Ala Tyr Ile Phe Asn Ser Asn Val Met Val Lys Val Ala Thr Gln Pro
 85 90 95
 Ala Asp Asn Pro Leu Asp Val Leu Ser Arg Lys Leu His Leu Gly Pro
 100 105 110
 Asn Val Gly Arg Asp Val Pro Arg Leu Ser Leu Pro Gly Lys Leu Val
 115 120 125
 Phe Pro Ser Ser Thr Gly Ser His Phe Ser Met Leu Gly Ile Gly Asp
 130 135 140
 Ile Val Met Pro Gly Leu Leu Leu Cys Phe Val Leu Arg Tyr Asp Asn
 145 150 155 160
 Tyr Lys Lys Gln Ala Ser Gly Asp Ser Cys Gly Ala Pro Gly Xaa Ala
 165 170 175
 Asn Ile Ser Gly Arg Met Gln Lys Val Ser Tyr Phe His Cys Thr Leu
 180 185 190
 Ile Gly Tyr Phe Val Gly Leu Leu Thr Ala Thr Val Ala Ser Arg Val
 195 200 205
 His Arg Ala Ala Gln Pro Ala Leu Leu Tyr Leu Val Pro Phe Thr Leu
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 Leu Pro Leu Leu Thr Met Ala Tyr Leu Lys Gly Asp Leu Arg Arg Met
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<210> 8

<211> 587
 <212> PRT
 <213> *Saccharomyces cerevisiae*

<400> 8

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Asn Lys Glu Leu Glu Gln Val Phe Glu Gln Ile Asn Ala Ile Val Glu
      35           40           45

Asn His Asn Asn Lys Leu Thr Thr Ala Phe Asp Lys Ile Ser Tyr Arg
      50           55           60

Val Ala His Lys Ile Thr His Leu Val Glu Ser His Ser Leu Val Phe
      65           70           75           80

Asn Tyr Ala Thr Leu Val Leu Ile Ala Ser Ala Leu Val Val Ile Gly
          85           90           95

Ser Phe Thr Ser Ile Ser Ser Ile Pro Phe Thr Ala Leu Pro Pro Thr
      100           105           110

Arg Glu His Ser Leu Phe Asp Pro Thr Asp Phe Asp Val Asp His Asp
      115           120           125

Cys His Val Ile Tyr Arg Glu Asn Asp Glu Asp Lys Lys Lys Lys Lys
      130           135           140

Lys Ser Lys Arg Phe Phe Asp Met Met Asp Glu Lys His Ala Ile Ile
      145           150           155           160

Leu Pro Leu Thr Ser Gly Cys Thr Leu Leu Ala Leu Tyr Phe Val Ile
          165           170           175

Lys Lys Leu His Leu Asn Trp Leu Lys Tyr Val Val Lys Ile Leu Asn
      180           185           190

Phe Asn Ile Thr Leu Leu Asn Ile Pro Ala Gly Thr Phe Val Tyr Ser
      195           200           205

Tyr Phe Leu Asn Ser Leu Phe Arg Asn Leu Ser His Leu Ala Ser Trp
      210           215           220

Asn Pro Leu Val Val Leu Pro Arg Tyr Arg Val Thr Ile Ala Asp Asp
      225           230           235           240

Asn Glu Asp Leu Asn Lys Ile Gly Gly Phe Val Thr Asn Leu Asn Tyr
          245           250           255

Lys Asp Gly Leu Thr Asn Ser Val Val His Lys Lys Thr Leu Asp Glu
      260           265           270

Ile Glu Lys Asp His Trp Met Lys His Phe Tyr Arg Arg Glu Leu Val
      275           280           285

Glu Pro Lys Asp Ile Lys Ser Lys Arg Gln Ile Ser Asn Met Tyr Leu
      290           295           300

Asn Ser Ala Leu Ile Val Ser Phe Val Leu Ser Ile Val Ser Thr Val
      305           310           315           320

Tyr Phe Tyr Leu Ser Pro Asn Asp Trp Leu Ile Ser Asn Ala Val Ser

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Phe Val Phe Gly Thr Asp Val Met Val Thr Val	Ala Thr Asn Leu Asp	
370	375	380
Ile Pro Val Lys Leu Ser Leu Pro Val Lys	Phe Asn Thr Ala Gln Asn	
385	390	400
Asn Phe Asn Phe Ser Ile Leu Gly Leu Gly Asp	Ile Ala Leu Pro Gly	
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Met Phe Ile Ala Met Cys Tyr Lys Tyr Asp Ile	Trp Lys Trp His Leu	
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Asp His Asp Asp Thr Glu Phe His Phe Leu Asn	Trp Ser Tyr Val Gly	
	435	440
Lys Tyr Phe Ile Thr Ala Met Val Ser Tyr Val	Ala Ser Leu Val Ser	
	450	455
Ala Met Val Ser Leu Ser Ile Phe Asn Thr	Ala Gln Pro Ala Leu Leu	
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Tyr Ile Val Pro Ser Leu Leu Ile Ser Thr Ile	Leu Val Ala Cys Trp	
	485	490
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	500	505
Val Asp Lys Ser Leu Lys Lys Ala Ile Glu Lys	Lys Glu Asn Ser Ile	
	515	520
Thr Tyr Ser Thr Phe Ile Leu Ser Glu Tyr Tyr	Asn Asp Ala Asp Lys	
	530	535
Tyr Ala Leu Leu Gly Asp Asp Val Asn Glu Asn	Phe Asp Asp Asp Glu	
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<211> 1776

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 9

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<210> 10
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 <212> DNA
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<210> 11
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 <213> Homo sapiens

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agcactacgc	ggccgccttc	cacgcccag	ggcatcgcg	tggcctacgg	cagcctcctg	120

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ctcatggcgc tgctgcccac cttcttcggc gccctgcgct ccgtacgctg cgcccgcggc 180
aagaatgctt cagacatgcc tgaacaatc accagccggg atgccgcccg cttccccatc 240
atcgccagct gcacactctt ggggctctac ctctttttca aaatattctc ccaggagtag 300
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aaggacctgg tgtgcctggg cctgagcagc atcggttggc tctggtacct gctgagggaag 540
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cacctcaaca atgtcagcac tggctgcac ctgctgggag gactcttcat ctacgatgtc 660
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acagagatgt tcagttatga ggagtcaaat cctaaggatc cagcggcagt gacagaatcc 1080
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<210> 12
<211> 1152
<212> DNA
<213> Homo sapiens

```

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<400> 12
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gaaaatcaag ataaggagaa agacagtaat agttcttctg ggtctttcaa tggcaacagc 180
accaataata gcatccaaac aattgactct acccaggctc tgttccttcc aattggagca 240
tctgtctctc ttttagtaat gttcttcttc tttgactcag ttcaagtagt ttttacaata 300
tgtacagcag ttcttgcaac gatagctttt gcttttcttc tcttcccgat gtgccagtat 360
ttaacaagac cctgctcacc tcagaacaag atttcctttg gttgctgtgg acgttttact 420
gctgctgagt tgctgtcatt ctctctgtct gtcatgctcg tcttcactcg gggttctact 480
ggccattggc ttctcatgga tgcactggcc atgggcctct gtgtcgccat gatcgccctt 540
gtccgcctgc cgagcctcaa ggtctcctgc ctgcttctct cagggcttct catctatgat 600
gtcttttggg tatttttctc agcctacatc ttcaatagca acgtcatggt gaaggtggcc 660
actcagccgg ctgacaatcc ccttgacgtt ctatcccgga agctccacct ggggcccact 720
gttgggcgtg atgttctctg cctgtctctg cctggaaaac tgggtcttccc aagctccact 780
ggcagccact tctccatggt gggcatcgga gacatcgta tgccctggct cctactatgc 840
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cccgccttc tctatttggg gccatttact ttattgccac tcttcacgat ggcctattta 1080
aaggcgacc tccggcggt gtggtctgag ccttttccact ccaagtccag cagctcccga 1140
ttcctggaag ta 1152

```

```

<210> 13
<211> 339
<212> DNA
<213> Mus musculus

```

```

<400> 13
gtattgggtt tcggagatat cattgtacca ggctgttga ttgcatattg tagaagattc 60
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atgatcatta cctttgttgt cctgatgggt atgaaaacag ggcagcctgc tctcctctac 180
ttggtacctt gtacacttat tactgtctca gtcgttgcct ggagtcgtaa ggaaatgaaa 240
aagttctgga aaggcagcag ctatcagggt atggaccacc tggactattc aacaaatgaa 300
gaaaatccag tgacgactga tgagcagatt gtacaacag 339

```

```

<210> 14
<211> 1134
<212> DNA
<213> Mus musculus

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<400> 14
atggattcgg ctgtcagcga tccgcacaac ggcagcgccg aggctggcac cccagccaac 60
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ctcatggcgc tgctgcccac cttcttcggc gccctgccgt cggtgcgctg cgcccgcggc 180
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cactggattg ccaacaacct gtttggcctg gccttctccc ttaatggggt agagctcctg 600
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ctgggacttg gagacattgt cattccaggg atcttcattg ccttactgct tcgttttgac 840
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gccgagatgt tcagttatga ggagtccaac cctaaagatc cagcagccgt gactgaatcc 1080
aaagaggagt caacagaggc gtcggcatcg aagaggctag agaagaagga gaaa 1134

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```

<210> 15
<211> 771
<212> DNA
<213> Mus musculus

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```

<220>
<221> misc_feature
<222> (524)..(525)
<223> n is a, c, g, or t

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<400> 15
cagaacaaga tttccttcgg ttgctgtggg cgttttactg ctgccgagct gctgtcgttc 60
tacctgtctg tcatgctcgt cctcatctgg gttctcactg gccactggct tctcatggat 120
gctctggcca tgggtctctg tgttgccatg atcgcccttc tccgcctgcc aagcctcaag 180
gtttcctgcc tgcttctctc agggcttctc atctacgat tcttctgggt gttcttctca 240
gcctacatct tcaacagtaa tgtcatgggt aaagtggcca cacagccagc tgacaatccc 300
ctcgacgttc tgtccaggaa gctccacctg ggacccaatg tggggcgtga tgttctctgc 360
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ggcatcgggg acatttgtgat gcccggcctc ctgttatgct ttgttcttcg ctatgacaac 480
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cgcctgcaga aggtctccta cttccactgc accctcatcg ggtactttgt aggtctgctc 600
actgcgactg tggcgtctcg cgtccaccga gctgccagc cagctctcct ctacttggtg 660
ccatttacc tattgccact cctcaccatg gcctacctaa agggtgactt acggaggatg 720
tggtctgagc ccttccactc caagtccagc agctcccgtt tcctggaagt a 771

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```

<210> 16
<211> 1761
<212> DNA
<213> Saccharomyces cerevisiae

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```

<400> 16
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gaacagatca atgctatagt tgaaaaccac aataacaaat taaccactgc ctttgataag 180
atatcatatc gcgttgctca caagattaca cacttgggtg aaagccattc tttagtattc 240
aactacgcca ctttagttct catcgcaagt gctttggctg ttattggctc atttacgtct 300
atttcttcta ttccatttac agctctacct cctacgagag aacactcatt gtttgatcct 360
acagattttg atgtggacca cgactgtcat gttatctacc gcgagaatga cgaagataaa 420
aagaaaaaga agaaaagcaa gaggtttttc gatatgatgg atgaaaaaca tgcgattata 480
ctgcccttaa ctagtggctg tactttactg gctctctatt ttgtgatcaa gaaactacac 540
ctaaactggc taaaatatgt ggtgaaaatt ttgaatttta atataacact gctaaatata 600
ccagctggca catttgtcta ctctacttt ctcaactcac ttttcagaaa cctatcacat 660

```

```

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accaactcag ttgttcacaa gaaaacattg gatgagattg aaaaagatca ttggatgaag 840
cattttttaca gaagagaatt agttgaaccg aaggatatta aatcgaagag gcagatcagc 900
aacatgtatt tgaatagcgc attaatgttt tcgttcgttc tgtccatcgt ttctaccgta 960
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atagaaaaaa aggagaactc tataacttat tcgaccttca tcctatcaga atactataat 1620
gatgccgaca agtatgcctt gcttgggtgat gatgtaaacg aaaattttga cgatgatgaa 1680
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ctcttggtat acgaatcttc t 1761

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<210> 17
 <211> 1560
 <212> DNA
 <213> Homo sapiens

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<400> 17
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tccattagtt tgatgaatct gacttccaca ccactatgca accttctga tattcctcct 240
gttggcataa agagcaaaagc agttgtgggt ccatggggaa gctgccattt tcttgaaaaa 300
gccagaattg cacagaaagg aggtgctgaa gcaatgttag ttgtcaataa cagtgtccta 360
tttctccctc caggtaacag atctgaattt cctgatgtga aaatactgat tgcatttata 420
agctacaaag acttttagaga tatgaaccag actctaggag ataacattac tgtgaaaatg 480
tattctccat cgtggcctaa ctatgattat actatgggtg gtatttttgg aattgcggtg 540
ttcactgggg cattaaagtgg atactggagt ggactagtgt aattggaaaa cttgaaagca 600
gtgacaactg aagatagaga aatgaggaaa aagaaggaag aatatttaac ttttagtcct 660
cttacagttg taatatttgt ggtcatctgc tgtgttatga tggcttact ttatttcttc 720
tacaaatggt tgggtttatgt tatgatagca attttctgca tagcatcagc aatgagtctg 780
tacaactgtc ttgctgcact aattcataag ataccatag gacaatgcac gattgcatgt 840
cgtggcaaaa acatggaagt gagacttatt tttctctctg gactgtgcat agcagtagct 900
gttgtttggg ctgtgtttcg aaatgaagac aggtgggctt ggattttaca ggatatcttg 960
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atcacaaaga atgggtgagag tatcatgggt gaactcgag ctggaccttt tggaaataat 1140
gaaaagtgtc cagtagtcat cagagtacca aaactgatct atttctcagt aatgagtgtg 1200
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gggcaacctg ctctcctcta tttagtacct tgcacactta ttactgcctc agttgttgcc 1440
tggagacgta aggaaatgaa aaagtctctg aaaggtaaca gctatcagat gatggaccat 1500
ttggattgtg caacaaatga agaaaaccct gtgatatctg gtgaacagat tgtccagcaa 1560

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<210> 18
 <211> 520
 <212> PRT
 <213> Homo sapiens

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<400> 18
Met Gly Pro Gln Arg Leu Ser Pro Ala Gly Ala Ala Leu Leu Trp
  1           5           10          15
Gly Phe Leu Leu Gln Leu Thr Ala Ala Gln Glu Ala Ile Leu His Ala
          20          25          30

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Ser	Gly	Asn	Gly	Thr	Thr	Lys	Asp	Tyr	Cys	Met	Leu	Tyr	Asn	Pro	Tyr	35	40	45
Trp	Thr	Ala	Leu	Pro	Ser	Thr	Leu	Glu	Asn	Ala	Thr	Ser	Ile	Ser	Leu	50	55	60
Met	Asn	Leu	Thr	Ser	Thr	Pro	Leu	Cys	Asn	Leu	Ser	Asp	Ile	Pro	Pro	65	70	75
Val	Gly	Ile	Lys	Ser	Lys	Ala	Val	Val	Val	Pro	Trp	Gly	Ser	Cys	His	85	90	95
Phe	Leu	Glu	Lys	Ala	Arg	Ile	Ala	Gln	Lys	Gly	Gly	Ala	Glu	Ala	Met	100	105	110
Leu	Val	Val	Asn	Asn	Ser	Val	Leu	Phe	Pro	Pro	Ser	Gly	Asn	Arg	Ser	115	120	125
Glu	Phe	Pro	Asp	Val	Lys	Ile	Leu	Ile	Ala	Phe	Ile	Ser	Tyr	Lys	Asp	130	135	140
Phe	Arg	Asp	Met	Asn	Gln	Thr	Leu	Gly	Asp	Asn	Ile	Thr	Val	Lys	Met	145	150	155
Tyr	Ser	Pro	Ser	Trp	Pro	Asn	Tyr	Asp	Tyr	Thr	Met	Val	Gly	Ile	Phe	165	170	175
Gly	Ile	Ala	Val	Phe	Thr	Gly	Ala	Leu	Ser	Gly	Tyr	Trp	Ser	Gly	Leu	180	185	190
Val	Glu	Leu	Glu	Asn	Leu	Lys	Ala	Val	Thr	Thr	Glu	Asp	Arg	Glu	Met	195	200	205
Arg	Lys	Lys	Lys	Glu	Glu	Tyr	Leu	Thr	Phe	Ser	Pro	Leu	Thr	Val	Val	210	215	220
Ile	Phe	Val	Val	Ile	Cys	Cys	Val	Met	Met	Val	Leu	Leu	Tyr	Phe	Phe	225	230	235
Tyr	Lys	Trp	Leu	Val	Tyr	Val	Met	Ile	Ala	Ile	Phe	Cys	Ile	Ala	Ser	245	250	255
Ala	Met	Ser	Leu	Tyr	Asn	Cys	Leu	Ala	Ala	Leu	Ile	His	Lys	Ile	Pro	260	265	270
Tyr	Gly	Gln	Cys	Thr	Ile	Ala	Cys	Arg	Gly	Lys	Asn	Met	Glu	Val	Arg	275	280	285
Leu	Ile	Phe	Leu	Ser	Gly	Leu	Cys	Ile	Ala	Val	Ala	Val	Val	Trp	Ala	290	295	300
Val	Phe	Arg	Asn	Glu	Asp	Arg	Trp	Ala	Trp	Ile	Leu	Gln	Asp	Ile	Leu	305	310	315
Gly	Ile	Ala	Phe	Cys	Leu	Asn	Leu	Ile	Lys	Thr	Leu	Lys	Leu	Pro	Asn	325	330	335
Phe	Lys	Ser	Cys	Val	Ile	Leu	Leu	Gly	Leu	Leu	Leu	Leu	Tyr	Asp	Val	340	345	350
Phe	Phe	Val	Phe	Ile	Thr	Pro	Phe	Ile	Thr	Lys	Asn	Gly	Glu	Ser	Ile	355	360	365
Met	Val	Glu	Leu	Ala	Ala	Gly	Pro	Phe	Gly	Asn	Asn	Glu	Lys	Leu	Pro	370	375	380

Val Val Ile Arg Val Pro Lys Leu Ile Tyr Phe Ser Val Met Ser Val
 385 390 395 400
 Cys Leu Met Pro Val Ser Ile Leu Gly Phe Gly Asp Ile Ile Val Pro
 405 410 415
 Gly Leu Leu Ile Ala Tyr Cys Arg Arg Phe Asp Val Gln Thr Gly Ser
 420 425 430
 Ser Tyr Ile Tyr Tyr Val Ser Ser Thr Val Ala Tyr Ala Ile Gly Met
 435 440 445
 Ile Leu Thr Phe Val Val Leu Val Leu Met Lys Lys Gly Gln Pro Ala
 450 455 460
 Leu Leu Tyr Leu Val Pro Cys Thr Leu Ile Thr Ala Ser Val Val Ala
 465 470 475 480
 Trp Arg Arg Lys Glu Met Lys Lys Phe Trp Lys Gly Asn Ser Tyr Gln
 485 490 495
 Met Met Asp His Leu Asp Cys Ala Thr Asn Glu Glu Asn Pro Val Ile
 500 505 510
 Ser Gly Glu Gln Ile Val Gln Gln
 515 520

<210> 19
 <211> 684
 <212> PRT
 <213> Homo sapiens

<400> 19
 Met Ala Cys Leu Gly Phe Leu Leu Pro Val Gly Phe Leu Leu Leu Ile
 1 5 10 15
 Ser Thr Val Ala Gly Gly Lys Tyr Gly Val Ala His Val Val Ser Glu
 20 25 30
 Asn Trp Ser Lys Asp Tyr Cys Ile Leu Phe Ser Ser Asp Tyr Ile Thr
 35 40 45
 Leu Pro Arg Asp Leu His His Ala Pro Leu Leu Pro Leu Tyr Asp Gly
 50 55 60
 Thr Lys Ala Pro Trp Cys Pro Gly Glu Asp Ser Pro His Gln Ala Gln
 65 70 75 80
 Leu Arg Ser Pro Ser Gln Arg Pro Leu Arg Gln Thr Thr Ala Met Val
 85 90 95
 Met Arg Gly Asn Cys Ser Phe His Thr Lys Gly Trp Leu Ala Gln Gly
 100 105 110
 Gln Gly Ala His Gly Leu Leu Ile Val Ser Arg Val Ser Asp Gln Gln
 115 120 125
 Cys Ser Asp Thr Thr Leu Ala Pro Gln Asp Pro Arg Gln Pro Leu Ala
 130 135 140
 Asp Leu Thr Ile Pro Val Ala Met Leu His Tyr Ala Asp Met Leu Asp
 145 150 155 160

Ile Leu Ser His Thr Arg Gly Glu Ala Val Val Arg Val Ala Met Tyr
 165 170 175
 Ala Pro Pro Glu Pro Ile Ile Asp Tyr Asn Met Leu Val Ile Phe Ile
 180 185 190
 Leu Ala Val Gly Thr Val Ala Ala Gly Gly Tyr Trp Ala Gly Leu Thr
 195 200 205
 Glu Ala Asn Arg Leu Gln Arg Arg Arg Ala Arg Arg Gly Gly Gly Ser
 210 215 220
 Gly Gly His His Gln Leu Gln Glu Ala Ala Ala Glu Gly Ala Gln
 225 230 235 240
 Lys Glu Asp Asn Glu Asp Ile Pro Val Asp Phe Thr Pro Ala Met Thr
 245 250 255
 Gly Val Val Val Thr Leu Ser Cys Ser Leu Met Leu Leu Leu Tyr Phe
 260 265 270
 Phe Tyr Asp His Phe Val Tyr Val Thr Ile Gly Ile Phe Gly Leu Gly
 275 280 285
 Ala Gly Ile Gly Leu Tyr Ser Cys Leu Ser Pro Leu Val Cys His Leu
 290 295 300
 Ser Leu Arg Gln Tyr Gln Arg Pro Pro His Ser Leu Trp Ala Ser Leu
 305 310 315 320
 Pro Leu Pro Leu Leu Leu Leu Ala Ser Leu Cys Ala Thr Val Ile Ile
 325 330 335
 Phe Trp Val Ala Tyr Arg Asn Glu Asp Arg Trp Ala Trp Leu Leu Gln
 340 345 350
 Asp Thr Leu Gly Ile Ser Tyr Cys Leu Phe Val Leu His Arg Val Arg
 355 360 365
 Leu Pro Thr Leu Lys Asn Cys Ser Ser Phe Leu Leu Ala Leu Leu Ala
 370 375 380
 Phe Asp Val Phe Phe Val Phe Val Thr Pro Phe Phe Thr Lys Thr Gly
 385 390 395 400
 Glu Ser Ile Met Ala Gln Val Ala Leu Gly Pro Ala Glu Ser Ser Ser
 405 410 415
 His Glu Arg Leu Pro Met Val Leu Lys Val Pro Arg Leu Arg Val Ser
 420 425 430
 Ala Leu Thr Leu Cys Ser Gln Pro Phe Ser Ile Leu Gly Phe Gly Asp
 435 440 445
 Ile Val Val Pro Gly Phe Leu Val Ala Tyr Cys Cys Arg Phe Asp Val
 450 455 460
 Gln Val Cys Ser Arg Gln Ile Tyr Phe Val Ala Cys Thr Val Ala Tyr
 465 470 475 480
 Ala Val Gly Leu Leu Val Thr Phe Met Ala Met Val Leu Met Gln Met
 485 490 495
 Gly Gln Pro Ala Leu Leu Tyr Leu Val Ser Ser Thr Leu Leu Thr Ser
 500 505 510

Leu Ala Val Ala Ala Cys Arg Gln Glu Leu Ser Leu Phe Trp Thr Gly
 515 520 525
 Gln Gly Arg Ala Lys Met Cys Gly Leu Gly Cys Ala Pro Ser Ala Gly
 530 535 540
 Ser Arg Gln Lys Gln Glu Gly Ala Ala Asp Ala His Thr Ala Ser Thr
 545 550 555 560
 Leu Glu Arg Gly Thr Ser Arg Gly Ala Gly Asp Leu Asp Ser Asn Pro
 565 570 575
 Gly Glu Asp Thr Thr Glu Ile Val Thr Ile Ser Glu Asn Glu Ala Thr
 580 585 590
 Asn Pro Glu Asp Arg Ser Asp Ser Ser Glu Gly Trp Ser Asp Ala His
 595 600 605
 Leu Asp Pro Asn Glu Leu Pro Phe Ile Pro Pro Gly Ala Ser Glu Glu
 610 615 620
 Leu Met Pro Leu Met Pro Met Ala Met Leu Ile Pro Leu Met Pro Leu
 625 630 635 640
 Met Pro Pro Pro Ser Glu Leu Gly His Val His Ala Gln Ala Gln Ala
 645 650 655
 His Glu Thr Gly Leu Pro Trp Ala Gly Leu His Lys Arg Lys Gly Leu
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 Lys Val Arg Lys Ser Met Ser Thr Gln Ala Pro Leu
 675 680

<210> 20
 <211> 2052
 <212> DNA
 <213> Homo sapiens

<400> 20
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 ctgttcagct ccgactacat caccctcccc cgggacctgc accacgcccc actcctgccc 180
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